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A Potential Blood Test for Transmissible Spongiform Encephalopathies by Detecting Carbohydrate-Dependent Aggregates of PrPres-Like Proteins in Scrapie-Infected Hamster Plasma

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Abstract: PrPres has rarely been detected in blood (except in leukocytes) even in diseased animal models that are known to contain a large amount of PrPres in infected tissues. It seems likely that PrPres detection in blood is difficult because of the low titer of infectious material within the blood. Here, we demonstrate the detection of proteinase K-resistant 3F4-reactive protein in the plasma of scrapie-infected hamsters but not in the plasma of mock-infected hamsters by partial purification using a novel method termed “acidic SDS precipitation,” in conjunction with a highly sensitive chemiluminescence detection system used to show the presence of PrP at a concentration equivalent to 1.4×10^{-9} g of brain homogenate or 1.5×10^{-12} g (6.5×10^{-17} mol) of rPrP by conventional Western blotting. The 3F4-reactive proteins in scrapie-infected hamster plasma often resulted in multiple Mw protein bands occurring at higher Mw positions than the position of the di-glycosyl PrP molecule. Mixing scrapie-infected hamster brain homogenate with mock-infected hamster plasma resulted in the formation of similar Mw positions for multiple 3F4-reactive proteins. Predigestion of carbohydrate side chains from the proteins in the plasma or brain homogenate before mixing resulted in failure to obtain these multiple 3F4-reactive proteins. These observations indicate that PrPres aggregated with other proteins in the plasma through carbohydrate side chains and was successfully detected in the plasma of scrapie-infected hamsters. Counterparts in these aggregates with PrPres-like proteins in scHaPI are not known but any that exist should resist the PK digestion.

Key words: PrPres-like protein, Carbohydrate, Scrapie infection, Discrimination

Transmissible spongiform encephalopathy (TSE) is a fatal infectious neurodegenerative disease. It is characterized pathologically by spongy deterioration of the central nervous system (CNS) and by the deposition of amyloid plaques composed of an abnormal isoform of the prion protein (PrP^{sc}) in infected tissues (1, 2, 19). An important biochemical property of PrP^{sc} is its partial resistance to protease digestion, which results in the formation of a β -sheet-rich isoform. This molecule has therefore also been called PrPres, and it has been considered a disease-specific entity associated with TSE (1, 2, 9). Although the vCJD epidemic in the U.K. is

declining, expansion of the disease throughout continental Europe and in many other countries has raised concern all over the world (9, 10, 28). After the appearance of three cases of transfusion-related vCJD infection

Abbreviations: 2× acidic saline, 0.02 M acetic acid containing 0.15 M NaCl and 10 mM EDTA-2Na; Brh, brain homogenates; CNS, central nervous system; KLH, keyhole limpet hemocyanin; mAb, monoclonal antibody; mc, mock-infected; Pl, plasma; PMCA, protein misfolding cyclic amplification; PrP^c, normal prion protein; PrPres, proteinase K-resistant prion protein; PrP^{sc}, disease-associated prion protein; PTA, phosphotungstic acid; PVDF, polyvinylidene fluoride; RES, reticuloendothelial system; rPrP, recombinant hamster PrP(25–233); SB, super block; sc, scrapie-infected; TBST, Tris Buffered Saline containing 0.05% Tw20; TSE, transmissible spongiform encephalopathy.

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in the U.K., transfusion-related iatrogenic expansion of vCJD between the asymptotically infected donor and blood recipients has caused growing concern (13, 21, 26). In addition, the U.K. government has recently reported a fourth case of vCJD associated with a blood transfusion (HPA Press statement; 18 Jan. 2007, abbreviated in 13). For this reason, the aim of research into developing an antemortem test has changed from detecting infected persons in an endemic area to estimating the population size of infected persons within a more global area in order to prevent the iatrogenic infection by tainted blood (4, 8, 28). The most useful tissues for the diagnostic confirmation of vCJD in humans are CNS and reticulo-endothelial system (RES) tissue as well as the tonsils and appendix (14–16). The tonsils and appendix have been used successfully for the histopathological detection of PrPres in epidemiological studies of vCJD infection in the U.K.: an extremely high frequency of infection was detected compared to the frequency of so-called classical CJD (16, 17, 22). However, it is difficult to sample the tonsils and the appendix in living subjects. Therefore, the pathological diagnosis of TSE is currently made principally on the basis of postmortem preparations of CNS tissues, highlighting the need for the development of a more rapid diagnostic method using body fluids, especially blood (6, 27). For this purpose, several methods have been proposed and examined for prophylactic use (23, 30, 32). However, none of these methods has proved to be sufficient for the purposes (5, 7, 20, 29). To achieve this goal, several problems must be solved; solutions include using preparations with minimally invasive sampling techniques and establishing an effective and specific method for detecting the disease marker with sufficient sensitivity (5). The first problem can be solved by using body fluids such as blood or urine as test specimens. Therefore, the key is to develop a system with sufficient sensitivity to detect PrPres in blood or urine (5, 6, 20, 22, 29). The presence of PrPres in the urine of TSE-infected animals and humans has been reported previously (33). However, it has been suggested that this uPrP^{sc} may be contaminated bacterial components in infected animal urine and not a marker of TSE (12). Blood has not been considered a highly infective source of classical CJD. The same was true for vCJD until the first victim of vCJD resulting from a blood transfusion was reported (21). This report was followed by reports of three more cases of possible transfusion-related transmission of vCJD (18, HPA Press statement; 18, Jan. 2007, 13). The development of a testing method using blood has therefore become a major goal of TSE research.

Here we show the successful detection (using a high-

ly sensitive chemiluminescence immunoblotting system) of a PrPres-like protein molecule in plasma collected from scrapie-infected (sc) but not from mock-infected (mc) hamsters. Although the infectivity of this molecule has not been tested, and the immunoblot pattern of the anti-PrP reactive protein in plasma (scHaPl) was somewhat different from that of the brain homogenate (scHaBrh) in scrapie-infected hamsters, the specific reactivity of these proteins to anti-PrP mAb, the demonstration of carbohydrate side chain-mediated association between PrPres and plasma proteins, and the removal of the carbohydrate chain resulted in the appearance of similar Mw proteins in scBrh and scPl firmly support the conjecture that the extra Mw proteins observed in the trial were the aggregates of PrPres and some plasma proteins.

Materials and Methods

Enzymes, monoclonal antibodies (mAb) and recombinant hamster PrP peptide. Proteinase K (PK: 40.0 mAnson units/mg protein) was purchased from Merck Co. (Rahway, N.J., U.S.A.). Peptide *N*-glycosidase F (PNGaseF, 25,000 units/mg protein) was purchased from Roche Diagnostics Co., Ltd.

The anti-PrP mAbs 3F4 (Signet, Mass., U.S.A.) and 6H4 (Prionics AG, Zürich, Switzerland) were stored in aliquots at -80°C until use. mAb 5C8-113 was prepared by immunizing PrP knockout mice with bovine recombinant PrP (Prionics AG); screening was conducted using the same molecule. TA180 and TA181 were provided by Medical Biological Laboratory (MBL) and were prepared by immunizing conventional Balb/c mice with synthetic peptides of the hamster PrP sequence CERYRE or CAVVGGLGGYML conjugated with keyhole limpet hemocyanin (KLH), respectively, then screened by the same peptides without KLH and conjugated with an ELISA plate. The epitope sites of the mAbs were 150–152 and 163–165 for TA180, and 129–131 for TA181. The epitope site of 5C8-113 has not yet been determined but is possibly an unknown conformation-dependent site. Anti-HIV P24 mAb (7A8.1; CHEMICON) was kindly donated by Dr. Iwakura of the Institute of Medical Sciences, Tokyo University and was used as a negative control for anti-PrP mAb reactions. Hamster recombinant PrP(25-233) (abbreviated rPrP hereafter) was purchased from Alicon AG (Switzerland).

Material from scrapie-infected and mock-infected hamsters. Twelve Syrian golden hamsters were inoculated with scrapie (Sc237)-infected hamster brain homogenate intra-cerebrally. Six hamsters were similarly inoculated with uninfected normal hamster brain

homogenate and were used as mock-infected hamsters. Hamsters from the two groups were anesthetized with ether at the terminal stage of disease among animals in the scrapie-infected group (approximately 50–70 days after inoculation) and after the same time interval among animals in the mock-infected group. Blood was collected from the animals with ACD containing 10 mM EDTA as an anticoagulant. Blood samples from scrapie-infected and mock-infected hamsters was centrifuged at low speed and the plasma fractions were collected (scPl and mcPl, respectively). Both scPl and mcPl were processed similarly thereafter. Brains were removed from the terminal-stage infected hamsters or the mock-infected hamsters and homogenized in TBS containing 0.5% NP40, 0.5% DOC and a protease inhibitor cocktail (Sigma) using a closed system homogenizer. These brain homogenates were then adjusted to a concentration of 10% with the above-mentioned buffer (scBrh^{crude} or mcBrh^{crude}, respectively). scBrh^{crude} or mcBrh^{crude} were centrifuged at low speed to remove insoluble materials, and the supernatant fractions (scBrh or mcBrh) were processed as described below.

Enzyme treatment. The plasma or brain homogenates were diluted 4-fold with TBS containing 10 mM EDTA and digested with PK (50 µg/ml) at 37 C for 60 min. These reactions were stopped by adding 1 mM Pefablock. The digestion step was omitted in a set of controls. The samples were then treated with 3% SDS and 50 mM DTT in TBS before being inactivated at 100 C for 10 min and stored at -80 C in small aliquots.

Acidic SDS precipitation. Stored preparations were inoculated with equal volumes of 0.02 M acetic acid containing 0.15 M NaCl and 10 mM EDTA-2Na (2× acidic saline) at 10 C, followed by centrifugation at 15,000 rpm for 10 min. The resulting precipitates were dispersed in Tris Buffered Saline (TBS) with 5 mM EDTA and inoculated with equal amounts of 2× acidic saline again. After further centrifugation, the resulting precipitates were rinsed with a 5-fold volume of methanol, then dissolved in Laemmli's SDS sample buffer and analyzed thereafter.

Immunoblot detection of PrP-like proteins. SDS-PAGE was carried out on a 15% gel using Laemmli's conventional buffer system. The electrophoresed proteins were transferred onto polyvinylidene fluoride (PVDF) membranes using a semidry system. After Western blotting, the membranes were blocked with Super Block (SB; PIERCE, Rockford, Ill., U.S.A.) for 1 hr at room temperature then overnight at 4 C. The blocked membranes were first washed three times with TBS containing 0.05% Tween 20 (TBST), then incubated with an anti-PrP monoclonal antibody (mAb; 3F4,

6H4 or similar), in SB containing 10% Block Ace (Dainippon Pharmaceutical Co., Ltd.) and 0.01% BSA for 1 hr at room temperature then overnight at 4 C thereafter. For maximum detection of protein signals, the blotted membranes were incubated overnight at 4 C. After incubation, the membranes were washed five times with TBST, incubated with horseradish peroxidase-conjugated goat anti-mouse IgG (HRPGAM) in TBST containing 10% Block Ace and 0.1% BSA for 1 hr as a secondary antibody, washed five more times with TBST and incubated with a chemiluminescence substrate (Super Signal West Femto Maximum Sensitivity Substrates: SSWF; PIERCE). To obtain optimum chemiluminescence signals, HRPGAM was used at a concentration of 2 ng/ml according to the manufacturers instruction manual and chemiluminescence signals for antibody-reactive bands were detected using an LAS3000 image analyzer (Fuji Film, Tokyo).

Results

Sensitivity of Detection Systems (Fig. 1)

To determine the limits of the detection system, rPrP, 10% homogenates of sc- or mcBrh (crude or low-speed centrifugation supernatant) and PK predigested sc- or mcBrh were used. They were abbreviated as sc- or mcBrh^{crude}, sc- or mcBrh^{sup} and sc- or mc BrhPK₅₀, respectively. In these experiments, the amount of PrP was indicated as brain equivalent (panel A) or brain protein (panel B) to enable convenient comparison between the equivalence to brain amount and brain protein. Protein amounts in the brain were determined before PK digestion. In panel A, 1.5×10^{-12} g (6.5×10^{-17} mol) for rPrP and PrPres in 1.4×10^{-9} g brain equivalent were detected. About 1/3 (equivalent to 2.25×10^{-6} g brain protein) of the PrP molecule in scBrh, and none of those in mcBrh looked like the PK-resistant molecule (PrPres; panel B). PrPres in scBrh was shifted from 30–32 kDa and 27–28 kDa before PK treatment to 25 and 20 kDa positions after PK treatment, respectively. As the total amount of PrPres plus PrPc in scBrh looked 3-fold larger than the amount of PrP in mcBrh, synthesis of the PrP was enhanced by scrapie infection in hamsters.

Discrimination of Scrapie Infection from Mock Infection by Plasma

When scrapie-infected or mock-infected Brh and plasma were pretreated with PK and subjected to immunoblot analysis, sc and mcBrh were easily discriminated by the PK treatment but sc and mcPl were not discriminated by the enzyme treatment (panel A). In mcPl, similar 3F4-reactive proteins were also

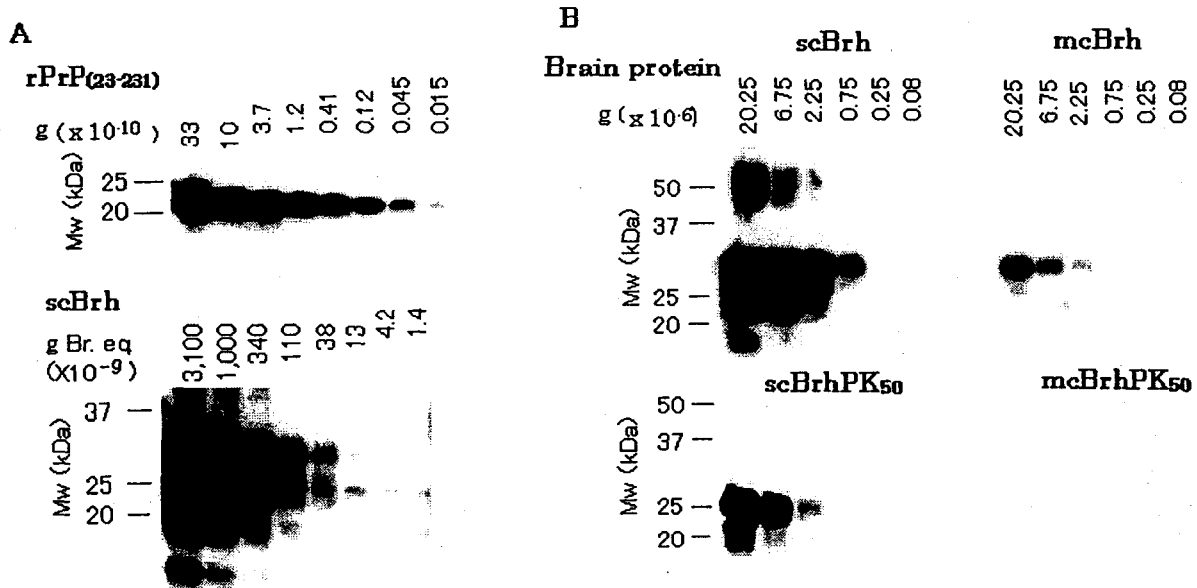


Fig. 1. Sensitivity and specificity determinations of immunoblot systems. Panel A: Sensitivity of used chemiluminescence immunoblots was determined using rPrP(25-233) (upper) or scBrh^{wt} (lower). These were diluted in the serial threefold manner as described in "Materials and Methods." Thereafter, each diluted preparation was subjected to chemiluminescence immunoblot detection. The amounts of each preparation used per lane are indicated in the figure as g ($\times 10^{-10}$) for rPrP(23-231) or g Br. eq. ($\times 10^{-9}$) for scBrh. Panel B: Specific detection of PrP in scBrh and mcBrh by chemiluminescence immunoblotting was indicated. scBrh or mcBrh (upper) or their PK-treated preparations (lower) were processed to serial threefold dilution series and subjected to chemiluminescence immunoblotting. PrP in each preparation was indicated per brain protein. Protein bands in scBrh showed the mixture of PrPres and PrPc. 3F4 and HRPgAM were used as the primary and secondary antibodies, respectively.

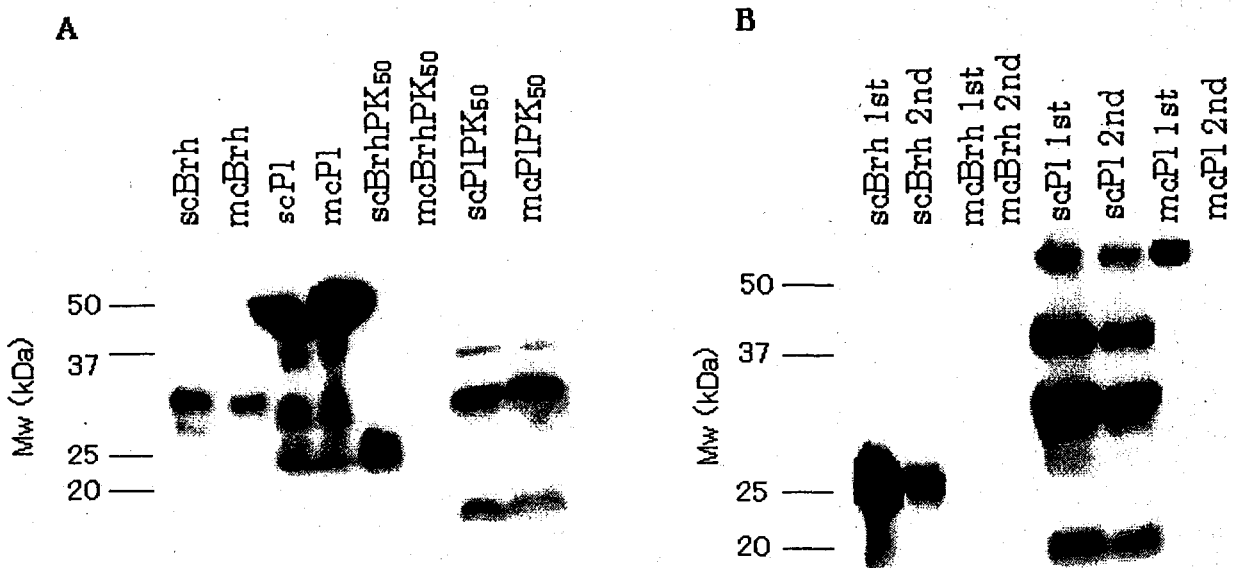


Fig. 2. Acidic SDS precipitation to discriminate scrapie infection and mock infection by their plasma. Panel A: scBrh, mcBrh, scPl and mcPl were processed and subjected to or not subjected to PK treatment. A chemiluminescence immunoblot analysis of PrP-like proteins was then performed using 3F4 primary and HRPgAM secondary antibodies. The PK treatment discriminated scBrh and mcBrh but did not discriminate between scPl and mcPl. Panel B: PK-treated scBrh, mcBrh, scPl and mcPl were subjected to acidic SDS precipitation condition, then analyzed by chemiluminescence immunoblotting. Acidic SDS precipitation condition was repeated twice (indicated as 1st and 2nd in the panel). This acidic SDS precipitation clearly discriminated scPl and mcPl as well as scBrh and mcBrh. The high MW protein band observed in the 1st precipitated fraction of mcPl was diminished in the 2nd precipitation. 3F4 and HRPgAM were used as the primary and secondary antibodies, respectively.

observed as in scPI. This observation was extremely different between the brain homogenate and plasma. From this observation, the presence of PK-resistant PrP molecules (PrPres-like molecules) in both sc- and mcPI was suspected (Fig. 2, panel A). These PrPres-like molecules in both plasma types have multiple inconstant Mw in experiments. Two patterns were often observed in 3F4-dependent immunoblot analysis; one was a 32 kDa major band with a 20 kDa minor band as well as 18, 25, 37 kDa faint bands (panel A; lane 7, 8), and the other was 20, 32, 40 and greater than 50 kDa dense bands as well as 27 kDa faint bands (Fig. 2, panel B; lane 5, 6).

The ability of the test to discriminate between scPI and mcPI was examined by acidic SDS precipitation (Fig. 2, panel B). PK-treated scBrh and mcBrh as well as scPI and mcPI preparations were subjected to acidic SDS precipitation condition (acidic SDS ppt) and analyzed with our immunoblotting system. In this experiment, precipitation was performed twice to ensure maximum removal of SDS soluble proteins. By this procedure, scPI and mcPI were clearly discriminated in the first precipitation and the higher Mw band that remained after the first precipitation was removed almost completely by the second precipitation. Discrimination between scPI and mcPI by acidic SDS precipitation was further confirmed by an experiment using 12 scPI and 6 mcPI samples (Fig. 3), but precipitation was only carried out once in this experiment. As shown in this figure, all 12 scPI samples showed the 3F4-reactive proteins but 4 mcPI samples did not. The mcPI of No. 1 and No. 6 showed weak 3F4-reactive bands. These observations confirm that scPI and mcPI can be

successfully discriminated using the acidic SDS precipitation but that precipitation should be repeated twice. Weak bands observed in the mcPI No. 1 and No. 6 were expected to disappear by performing one more acidic SDS precipitation procedure.

Effect of Deglycosylation

It is known that three Mw species of PrP, di-, mono- and none-glycosylated molecules, exist in the brain and deglycosylation of the molecules causes the three protein species to accumulate into a single Mw. So, in order to determine whether deglycosylation affects the formation of multiple Mw protein bands in sc or mcPI, PK-treated sc and mcPI were deglycosylated or further processed using the acidic SDS precipitation procedure then compared to similarly processed scBrh. As shown in Fig. 4, 20–27 kDa proteins in scBrh and 19–50 kDa multiple Mw proteins in scPI and mcPI were detected following PK treatment (step 1). With deglycosylation of scBrh by PNGase F treatment, large amounts of 18 kDa protein appeared as was expected. Deglycosylation of scPI and mcPI resulted in 18 kDa proteins appearing but multiple higher Mw protein bands remained (step 2). After acidic SDS precipitation of these PK digested and deglycosylated materials, the multiple higher Mw protein bands in scPI disappeared, whereas a small amount of discrete 18 kDa protein bands remained in scBrh and scPI. These protein bands were not detected following similar treatment of mcPI (step 3). A long period of exposure (10 min) was necessary to obtain the protein signals described from step 3 of the experiment because the PrP-like proteins were difficult to detect after the deglycosylation step of the

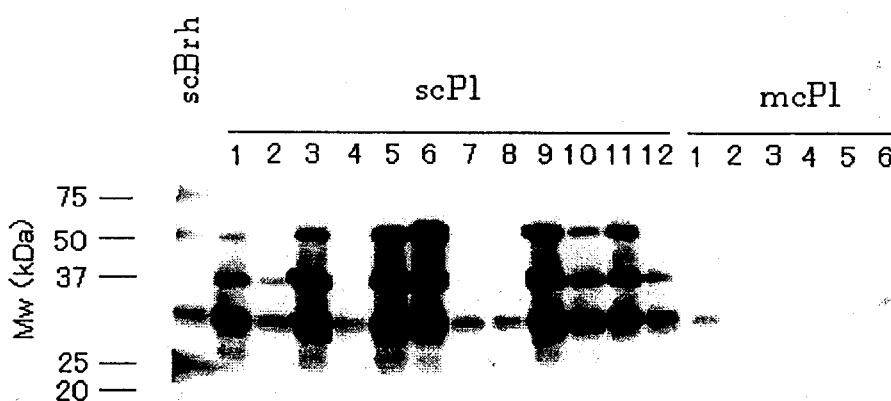


Fig. 3. Discrimination of PK-treated scPI and mcPI by acidic precipitation. Twelve preparations of scPI and 6 preparations of mcPI were pretreated with PK (50 μ g/ml). scBrh was similarly treated before processing. After the pretreatment, the scPI and mcPI as well as scBrh were processed to the acidic SDS precipitation stage and analyzed by the normal immunoblot systems as described in "Materials and Methods." Anti-PrP mAb 3F4 and HRP-GAM were used as the primary and secondary antibodies, respectively, for the immunoblot analysis.

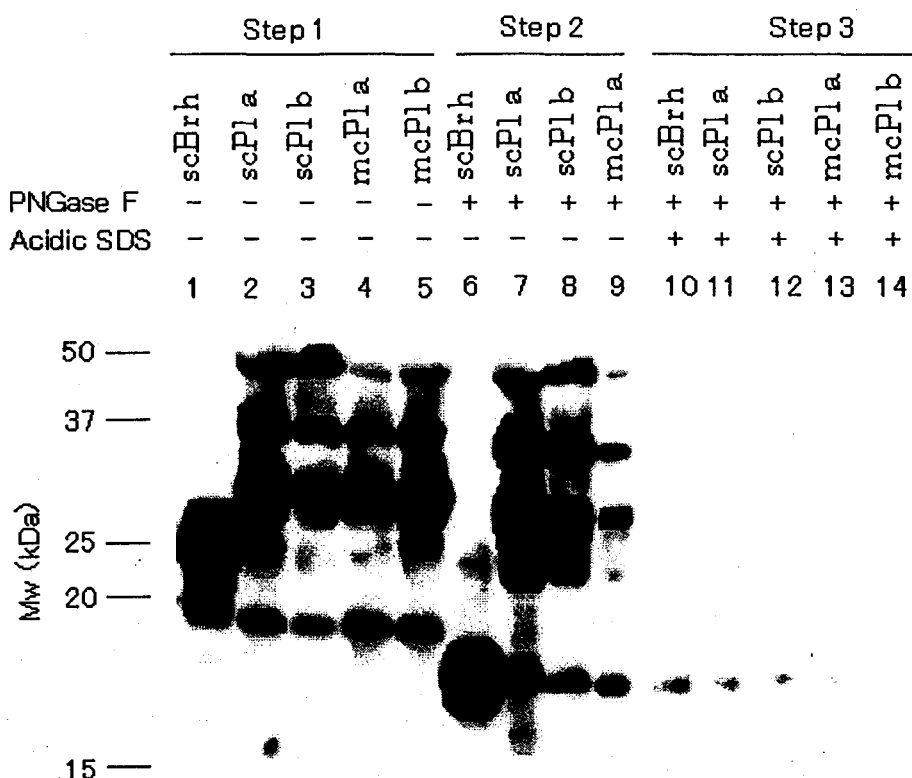


Fig. 4. Disappearance of extra Mw protein bands by digestion of carbohydrate side chains followed by acidic SDS precipitation. scBrh, two preparations of scPl and two preparations of mcPl were treated with PK (step 1, lanes 1–5). The PK-treated preparations were digested by PNGase F to remove the carbohydrate side chain on the protein molecules (step 2, lanes 6–9) then processed finally to the acidic SDS precipitation stage (step 3, lanes 10–14). Lanes were: Brain homogenate: 1, 6, 10; scPl: 2, 3, 7, 8, 11, 12; and mcPl: 4, 5, 9, 13, 14. The immunoblot pattern of each preparation during the three steps was determined. For the immunoblot analysis, 3F4 mAb and HRP-GAM were used as the primary and secondary antibodies, respectively. Preparations treated with PNGase F or acidic SDS precipitation are indicated as (+) and untreated or unprocessed preparations are indicated as (–) in the figure. 3F4 and HRP-GAM were used as the primary and secondary antibodies, respectively.

acidic SDS precipitation protocol. These observations suggested that carbohydrate side chains were involved in the formation of the multiple higher Mw protein bands (Fig. 4).

Appearance of Multiple Higher Mw Proteins by Mixing PK-Treated scBrh and PK-Treated or PK-Untreated mcPl

PK-treated Brh was mixed with PK-treated or PK-untreated mcPl then processed to the acidic SDS precipitation stage. These mixed preparations were compared with preparations of unmixed components in immunoblot analysis. Mixing of the PK-treated scBrh with PK-treated or PK-untreated mcPl resulted in the formation of higher Mw multiple protein bands as observed in scPl. Mixing with PK-treated mcPl seemed to show more discrete bands than mixing with PK-untreated mcPl. In PK-treated mcHaBrh, PK-treated or PK-untreated mcPl, these higher Mw protein bands

were not observed. These immunoblot results suggested that the multiple Mw 3F4-reactive proteins were newly formed by the association between PrPres in scBrh and some PK-resistant plasma proteins in mcPl (Fig. 5).

Effect of Deglycosylation for the Association of PrPres in scBrh and PK-Resistant Protein in Plasma

As the deglycosylation of scPl resulted in failure to form the multiple higher Mw proteins but resulted in the appearance of a discrete 18 kDa band. As the Mw of which is similarly to the deglycosylated PrPres in scBrh, the possible involvement of saccharide chains was suspected for the formation of multiple extra Mw protein bands. To confirm this possibility, PK-pretreated scBrh and mcPl were deglycosylated by PNGase F or left untouched. After mixing the two preparations in the combination indicated in Fig. 6, acidic SDS precipitation was performed thereafter. As 3% SDS in the stored plasma or brain homogenates inhibits deglyco-

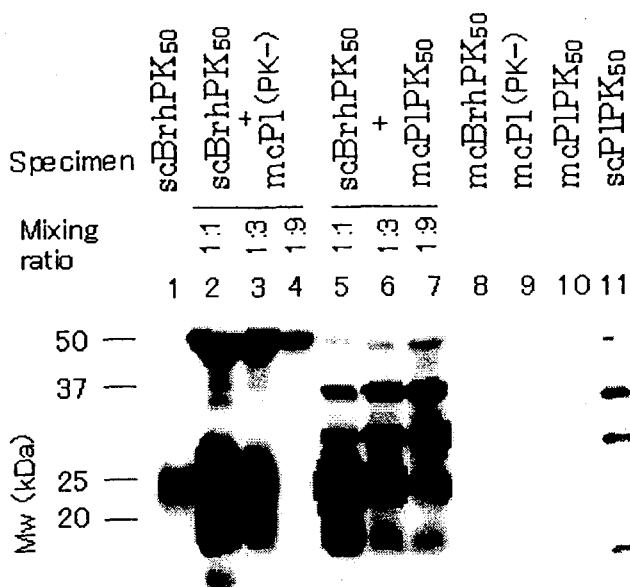


Fig. 5. Appearance of extra Mw proteins by mixing scBrh and mcPI after the acidic SDS precipitation. Proteinase K-pretreated scBrh was mixed with PK-treated or untreated mcPI and diluted to 1:3 or 1:9 in the presence of the PK-treated or untreated mcPI preparations. Then the mixed and unmixed preparations were processed to the acidic SDS precipitation stage. These processed preparations were compared by immunoblot analysis using 3F4 mAb and HRPgAM as the primary and secondary antibodies, respectively. Lanes: 1: PK treated scBrh; 2-4: PK-treated scBrh was mixed with an equal amount of PK-untreated mcPI (lane 2), diluted to 1:3 (lane 3), diluted to 1:9 (lane 4); 5-7: PK-treated scBrh was mixed with an equal amount of PK-treated mcPI (lane 5), diluted to 1:3 (lane 6), diluted to 1:9 (lane 7); 8: PK-treated mcBrh; 9: PK-untreated mcPI; 10: PK-treated mcPI; 11: PK-treated scPI. 3F4 and HRPgAM were used as the primary and secondary antibodies, respectively.

sylation reaction by PNGase F, the preparations for deglycosylation were diluted 30-fold before the reaction. After the deglycosylation, proteins in the reaction mixture of PNGase F treatment were precipitated by methanol and dissolved again to their original volumes with a primary buffer system that contained 3% SDS before mixing. Acidic SDS precipitation after the mixing of these deglycosylated preparations resulted in the appearance of an 18 kDa discrete band with a similar Mw to the deglycosylated PrPres in scBrh. Deglycosylation of brain proteins as well as of plasma proteins separately failed to form the higher Mw multiple protein bands. Mixing of PNGase F-treated scBrh and PNGase F-untreated mcPI formed a somewhat large amount of discrete 18 kDa proteins (lane 5).

Discussion

For antemortem diagnostic tests, body fluids such as

blood or urine may be the most convenient specimens. The infectivity of blood in TSE-infected animals has already been determined to be 10–30 ID₅₀/ml (5). For this reason, the detection sensitivity of PrPres in blood is required to be in the order of ng/ml to test for TSE as has been mentioned elsewhere. Immunoblotting systems cannot detect such a low level of PrPres even in the blood of experimentally infected animals, so more sensitive methods to detect lower concentrations of PrP molecules need to be developed for antemortem diagnostic tests using blood or other body fluids. Various trials by several investigators have attempted to solve this extremely difficult problem (7, 19, 20, 30). In these studies, capillary electrophoresis analysis using a fluorescence-labeled synthetic PrP peptide, a combination of conformation-dependent PTA precipitation and ELISA, PCR of synthetic RNA conjugated with anti-PrP mAb and *in vitro* multiplication of abnormal PrP isoform (Protein Misfolding Cyclic Amplification; PMCA) have been suggested (3, 20, 29, 30). The PMCA method was shown to detect the presence of PrPres in scrapie-infected pre-mortem hamster blood using the buffy coat lysate (29). However, because these methods are complex and require a long time to obtain final results, their use in blood screening may be restricted. On the other hand, the common immunoblotting system used after PK treatment is excellent for detecting PrPres in the CNS or in other disease-affected tissues of infected animals. However, the usual immunoblot detection is less sensitive than the methods mentioned above. Therefore, a method that uses the common immunoblotting system would be the first choice for an antemortem test if its detection sensitivity could be enormously enhanced. It is suspected that the detection of PrPres molecules in blood is made more difficult by contamination from a large amount of protein, and so a method that will selectively concentrate the PrPres in blood to allow detection is therefore required. We tried to use the common immunoblotting systems in combination with a selective concentration method for PrPres-like protein aggregates and a highly sensitive chemiluminescence method. Using this combination, we successfully showed the presence of PrPres-like proteins in the scPI by means of reactivity to several anti-PrP mAbs, and by the similarity of Mw with the PrPres in infected hamster brains after deglycosylation. Moreover, carbohydrate may cause the PrPres-plasma protein aggregation and form the multiple Mw 3F4-reactive PrP-like proteins. PrP is a membrane protein and is known to aggregate frequently, especially after conversion to its disease-associated abnormal isoform. For this reason, detection of these aggregates is also the optimal way to develop an assay

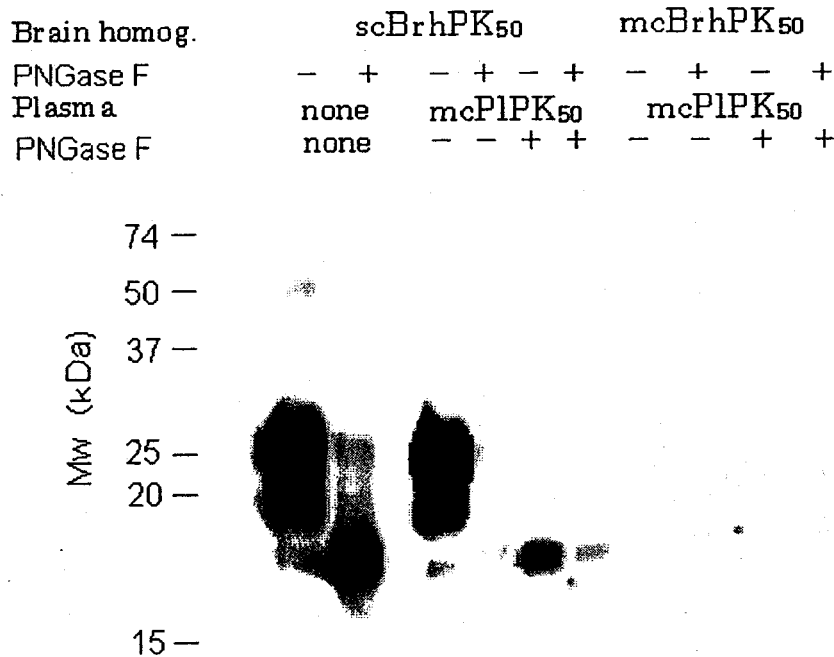


Fig. 6. Inability to form aggregate between scBrh and mcPl by digestion of carbohydrate before mixing. PK-treated scBrh, mcBrh and mcPl were further treated or not treated with PNGase F to digest the carbohydrate side chains on the proteins. These pretreated preparations were mixed with each other as indicated in the figure, and processed to the acidic SDS precipitation stage after mixing. Lanes: 1: scBrh (PNGase F-), 2: scBrh (PNGase F+); 3: scBrh (PNGase F-) mixed with mcPl (PNGase F-); 4: scBrh (PNGase F-) mixed with mcPl (PNGase F+); 5: scBrh (PNGase F+) mixed with mcPl (PNGase F-); 6: scBrh (PNGase F+) mixed with mcPl (PNGase F+); 7: mcBrh (PNGase F-) mixed with mcPl (PNGase F-); 8: mcBrh (PNGase F-) mixed with mcPl (PNGase F+); 9: mcBrh (PNGase F+) mixed with mcPl (PNGase F-); 10: mcBrh (PNGase F+) mixed with mcPl (PNGase F+), in which (PNGase F+) and (PNGase F-) mean digested or non-digested with PNGase F before mixing, respectively. 3F4 and HRP_{GAM} were used as the primary and secondary antibodies, respectively.

method when using blood. However, previous tests for evaluating the sensitivity of detection systems using PrP molecules have frequently failed, presumably due to the tendency of the PrP molecule to form aggregates. We therefore evaluated the sensitivity of the detection system using SDS sample buffer which contained 0.1% BSA for the dilution buffer and by boiling the preparation throughout the serial dilution steps. This method allowed us to obtain a proper dispersion of the PrP aggregate in the test preparation and we successfully showed that the endpoint of the detection system was 1.5×10^{-12} g (6.05×10^{-17} mol) or more of rPrP and PrPres in 1.4×10^{-9} g brain equivalent of scHaBrh. As the scBrh has an infectivity titer of 10^{-7} – 10^{-9} ID₅₀/ml, this chemiluminescence system can detect PrPres corresponding to 1 ID₅₀/ml or more, which is sufficiently greater than the value required to detect PrPres in blood (Fig. 1). We therefore decided to use this chemiluminescence system to detect PrPres in scPl. This system also allowed us to determine the detection limit of PrP protein in the brain (Fig. 1B).

Adding the acidic SDS precipitation stage to the pro-

cedure enabled successful discrimination of scPl and mcPl. The acidic SDS condition may selectively target aggregated PrP molecules, suggesting that PrP molecules in mcPl may not be aggregated. This observation is reasonable in that one of the main differences between PrPres and PrPc may be whether they exist in an aggregated form or not. Some investigators have tried to obtain PrPres in blood in an aggregated form (7). The aggregation of PrPres is thought to be a result of the more hydrophobic nature of the PrPres molecule than that of PrPc (24). However, the phenomenon observed here clearly suggests that an important factor for aggregation may be the presence of a carbohydrate side chain on both PrPres and plasma proteins rather than the hydrophobic nature of the PrPres. Carbohydrate has often been described as the outfilter for glycosylation and function (25).

The types of protein that aggregate with the PrPres-like molecules are not known. Some plasma proteins are known to associate with the PrP, but it is possible that the PrP molecules in hamster plasma may also be a candidate for these plasma proteins (11, 31, 34). PK-

resistant PrP molecules have recently been reported in uninfected human brains as well as in uninfected mouse and hamster brains and have been labeled a silent prion. PK-resistant protein in mcPI, which is able to aggregate with PrPres could be the silent prion in hamster plasma (34). Weakly observed 3F4-reactive protein bands in mcPI suggest the existence of the silent prion in plasma (Fig. 3).

In the lanes of scPI-2, -4, -7, -8, discrete bands in the Mw 32 kDa region were observed without other bands. As the band of this Mw region was weakly observed in mcPI-1 and scPI-6, it is somewhat difficult to decide the positive expression of 3F4-reactive protein for scPI-2, -4, -7 and -8. However, intensities of these signals in the scPI preparations were obviously strong compared to the signals in mcPI preparations. Thus it may be difficult to decide positive or not positive by performing acidic precipitation just one time. It is reasonable that an individual animal does not express the 3F4-reactive protein similarly in a time dependent manner and expression strength. Therefore, if blood testing is introduced, the plasma preparation should be processed twice with this acidic SDS precipitation, and the test should be conducted several times at different times.

Here we showed the successful discrimination of scrapie-infected and mock-infected hamsters by their plasma preparations using a novel combination method termed acidic SDS precipitation along with a highly sensitive chemiluminescence immunoblot system. In the immunoblots of PK-treated plasma preparations, multiple protein bands at Mw higher than the 25 kDa position were observed. These protein bands were observed in both scPI and mcPI after PK treatment. As these proteins were 3F4-reactive as well as PK-resistant, they were very likely to be PrPres molecules. However, observations showing multiple bands of higher than 25 kDa in Mw in mcPI as well as in scPI were very different from the electrophoresis pattern of scBrh. These differences between plasma and Brh have to be explained if the multiple PrPres-like proteins in plasma are aggregates of PrPres and some other plasma protein. This is similar with an observation in which the C-terminal domain of a recombinant mouse PrP peptide was aggregated spontaneously even in SDS sample buffer (24). Differences of electrophoresis patterns in Fig. 5, lanes 5-7 and Fig. 6, lane 3 or Fig. 2B, lane 5, 6 in spite of the same processing protocol may explain in which aggregation counterparts with PrPres in these plasma preparations may not be the same molecule, in preparation. After the PK treatment, an enormous amount of partial peptides was distributed in the broad Mw region if total protein was stained on WB membrane. This means that multiple partial peptides which

possessing carbohydrate chains may have the potential to become the counterpart of these aggregates. We could not control the combination of the molecules. A deglycosylation experiment using both scPI and scBrh solved this question. After deglycosylation and acidic SDS precipitation, both scBrh and scPI showed a single discrete protein band at the 18 kDa Mw position. This observation strongly suggests that the carbohydrate side chain might be an important factor in the aggregation of the PrPres-like protein with some other proteins. From these observations, one of the components required to form aggregates must be the PrPres molecule but the other component need not be another PrPres molecule. That is, both self aggregation as well as aggregation of multiple hetero molecules could be resulted in the formation of the multiple Mw protein bands. Although dense bands at 25 kDa was observed in the scBrh and mcPI mixing (Fig. 5, lanes 5-7), the bands were obscure in scPI (Fig. 2, lanes 5, 6). This discrepancies between the preparations may conjectured by the differences of PrPres and plasma protein ratio. In Fig. 5, lanes 5-7, larger amount of scBrh compared to mcPI showed pattern more similar to that of scBrh, larger amount of mcPI showed more discrete band pattern after the PK treatment, in reverse. In this observation, 20 and 25 kDa protein bands were decreasing gradually along with mcPI was increasing. Therefore, it is conjectured that the 20 and 25 kDa proteins were not observed if less amounts of PrPres existed in scPI as observed in Fig. 2B, lane 5, 6.

PrPres was also found in uninfected human brains and labeled a silent prion (34). Similar molecules are likely to be present in non-infected hamsters and in mouse brains as well. If the silent prion in hamster and mouse brains is also exist in plasma, the PK-resistant 3F4-reactive proteins observed in mcPI in this experiment may be the candidate in hamster plasma. The silent prions in hamster plasma could aggregate with themselves or with other proteins to form the multiple higher Mw proteins in mcPI as well as scPI. But if the silent prion exists in mcPI, it must be discriminated through the blood tests. The acidic SDS precipitation process reported here may be useful for such trials.

So, as the PrPres molecules in hamster, 25 kDa, 20 kDa and 18 kDa proteins correspond to the di-, mono and no carbohydrate molecules, respectively. Multiple higher Mw protein bands were presumably aggregates with PrPres and other plasma proteins. The phenomenon that Mw of these aggregates were not found within a constant range indicated that counterparts of presumable PrPres might not be the specialized molecules in preparations; the silent prion may be included within these inconsistent molecules. Furthermore, we could